

Figure 1

Sequence alignment of mouse Serca 1, 2 and 3 protein.

5	Serca1a	1	MEAAHSKSTEECLSYFGVSETTGLTPDQVKRHLEKYGNELPAEEGKSLWELVVEQFEDL
	Serca2a	1	..N..T.TV..V.GH...N.S...SLE...KLK.RW.S.....T.L...I.....
	Serca2b	1	..N..T.TV..V.GH...N.S...SLE...KLK.RW.S.....T.L...I.....
	Serca3a	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca3b	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
10	Serca3c	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca1a	61	LVRILLLLAACISFVLAWFEEGEETVTAFVEPFVILLILIANAIVGVWQERNAENAIEALK
	Serca2a	61I.....V.....
	Serca2b	61I.....V.....
15	Serca3a	61LV.....T.....L..M...V.....S.....
	Serca3b	61LV.....T.....L..M...V.....S.....
	Serca3c	61LV.....T.....L..M...V.....S.....
	Serca1a	121	EYEPENGKQYRADRKSVQRIKARDIVPGDIVEVAVGDKVPADIRILSIKSTTLRVDQSIL
20	Serca2a	121Q.....K.....I.....LT.....
	Serca2b	121Q.....K.....I.....LT.....
	Serca3a	121I.S...G...R.....L.LIE.....
	Serca3b	121I.S...G...R.....L.LIE.....
	Serca3c	121I.S...G...R.....L.LIE.....
25	Serca1a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAVGIVATTGVSTEIGKIRDQMA
	Serca2a	181M.V.VA...N.....E.V
	Serca2b	181M.V.VA...N.....E.V
	Serca3a	181T...AI.....S...L.VAVA..LQ..L...S...
30	Serca3b	181T...AI.....S...L.VAVA..LQ..L...S...
	Serca3c	181T...AI.....S...L.VAVA..LQ..L...S...
	Serca1a	241	ATEQDKTFLQQLDEFGEQLSKVISLICVAVWLINIGHFNDPVHGGSWFRGAIYYFKIAV
	Serca2a	241	...ER.....I...I.....I.....
35	Serca2b	241	...ER.....I...I.....I.....
	Serca3a	241	.V.PER...R.....R...HA..V.....V.....A..A....L...V.....
	Serca3b	241	.V.PER...R.....R...HA..V.....V.....A..A....L...V.....
	Serca3c	241	.V.PER...R.....R...HA..V.....V.....A..A....L...V.....
40	Serca1a	301	ALAVAAIPEGLPAVITTCLALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTNNQ
	Serca2a	301
	Serca2b	301
	Serca3a	301R.....
	Serca3b	301R.....
45	Serca3c	301R.....
	Serca1a	361	MSVCKMFIIDKVDGDVCSLNEFSITGSTYAPEGEVLKNDKPVRAGQYDGLVELATICALC
	Serca2a	361	...R...L...E..T.....I...Q.D...KCH.....
	Serca2b	361	...R...L...E..T.....I...Q.D...KCH.....
50	Serca3a	361	...R..VVAEAEAGT.R.H..T.S.T..T.....RQGEQ...C..F.....
	Serca3b	361	...R..VVAEAEAGT.R.H..T.S.T..T.....RQGEQ...C..F.....
	Serca3c	361	...R..VVAEAEAGT.R.H..T.S.T..T.....RQGEQ...C..F.....
	Serca1a	421	NDSSLDFNETKGVEKVGGEATETALTTLVEKMNVFNTEVRSLSKVERANACNSVIRQLMK
55	Serca2a	421	...A..Y..A.....C.....D..LKG...I.....K....
	Serca2b	421	...A..Y..A.....C.....D..LKG...I.....K....
	Serca3a	421	...A..Y..A.....C.....D..DLKG..R....G.....K...R
	Serca3b	421	...A..Y..A.....C.....D..DLKG..R....G.....K...R
	Serca3c	421	...A..Y..A.....C.....D..DLKG..R....G.....K...R

	Serca1a	481	KEFTLEFSRDRKSM SVYCSPAKSSRAAVGNKMFVKGAPEGVIDRCNYVRVGTTTRVPLTGP
	Serca2a	481T.N.P..TSMS-.....THI...S.K..M.PG
	Serca2b	481T.N.P..TSMS-.....THI...S.K..M.PG
	Serca3a	481T.TRADEPKVQ.S.....S.E..SS....SRTA..STT
5	Serca3b	481T.TRADEPKVQ.S.....S.E..SS....SRTA..STT
	Serca3c	481T.TRADEPKVQ.S.....S.E..SS....SRTA..STT
	Serca1a	541	VKEKIMSVIKEWGTGRDRTLRLALATRDTPPKREEMVLDDSAKFM EYEMDLTFVGVVGML
	Serca2a	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN....C....
10	Serca2b	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN....C....
	Serca3a	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3b	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3c	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
15	Serca1a	601	DPPRKEVTGSIQLCRDAGIRVIMITGDNKGTAIAICRRIGIFSENEEVTDRA YTGREFDD
	Serca2a	600I..AS.VK...Q.....V.....GQD.D..SK.F.....E
	Serca2b	600I..AS.VK...Q.....V.....GQD.D..SK.F.....E
	Serca3a	601P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
	Serca3b	601P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
20	Serca3c	601P..AAC.TR.SR....V.....V.....L...GDT.D.LGK.....
	Serca1a	661	LPLAEQREACRRACCFARVEPSHKSKI VEYLQSYDEITAMTGDGVNDAPALKKAEIGIAM
	Serca2a	660	.SPSA..D..LN.R.....F...F.....S.....
	Serca2b	660	.SPSA..D..LN.R.....F...F.....S.....
25	Serca3a	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca3b	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca3c	661	.SPEQ..Q...T.R.....A...R...N...FN.....
	Serca1a	721	GSGTAVAKTASEMVLADDNFSTIVA AVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAA
30	Serca2a	720
	Serca2b	720
	Serca3a	721S.A...S...AS.....I
	Serca3b	721S.A...S...AS.....I
	Serca3c	721S.A...S...AS.....I
35	Serca1a	781	LGLPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMDRPPRSPKEPLISGWLFFRYMAI
	Serca2a	780	..F.....NK...N.....L..
	Serca2b	780	..F.....NK...N.....L..
	Serca3a	781EK...N.R.A.....L..
40	Serca3b	781EK...N.R.A.....L..
	Serca3c	781EK...N.R.A.....L..
	Serca1a	841	GGYVGAATVGAAAWFLYAEDGPHVSYHQLTHFMQCTEHNPEFDGLDCEVF EAPEPMTMA
	Serca2a	840	.C.....IA.DG..R..FY..S..L..K.D..D...V..AI..S.Y.....
45	Serca2b	840	.C.....IA.DG..R..FY..S..L..K.D..D...V..AI..S.Y.....
	Serca3a	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
	Serca3b	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
	Serca3c	841	.V...L...A..T....DAE..Q.TFY..RN.LK.S.D..L.A.I..K...SRF.T...
50	Serca1a	901	LSVLVTIEMCNALNSLSENQSLLRMP PVWNIWLLGSICLSMSLHFLILYVDPLPMIFKLR
	Serca2a	900E...V.....E...L..QIT
	Serca2b	900E...V.....E...L..QIT
	Serca3a	901V.....L.P...AVVM..A.....L.P...L..QVT
	Serca3b	901V.....L.P...AVVM..A.....L.P...L..QVT
55	Serca3c	901V.....L.P...AVVM..A.....L.P...L..QVT

Serca1a 961 ALDFTQWLMVLKISLPVIGLDELLKFIARNYLEG
Serca2a 960 P.NL.....LM..T...V.....QPAILE
Serca2b 960 P.NL.....LM..T...V.....QPGKECVQPATKSSCSLSACTDGISWP
Serca3a 961 P.SGR..GV..QM.....L...A..YLS..HMDEKKDLK
5 Serca3b 961 P.SGR..GV..QM.....L...A..YLS..HMD.VLGTFMQARSRLPTTSRTPYHTGKK
Serca3c 961 P.SGR..GV..QM.....L...A..YLS..HMD.VLGTFMQARSRLPTTSRTPYHTGLA

Serca2b 1020 FVLLIMPLVVVVYSTDTNFSDFMFW
10 Serca3b 1021 GPEVNPGSRGESPVWPSD
Serca3c 1021 SWKKRT

Figure 2 **Sequence similarity of Serca2 proteins in mammalian species**

	Mouse_2a	1	MENAHKTKTVEEVLGHFGVNESTGLSLEQVKLKERWGSNELPAEEGKTLLELVIEQFEDL
5	Mouse_2b	1
	Rat_2b	1
	Rat_2a	1
	Dog_2a	1
	Cat_2a	1Y.....
10	Pig_2a	1
	Pig_2b	1
	Human_2b	1
	Human_2c	1
	Human_2a	1
15	Rabbit_2a	1
	Rabbit_2b	1
	Mouse_2a	61	LVRILLLLAACISFVLAWFEEGEETITAFVEPFVILLILVANAIVGVWQERNAENAIEALK
	Mouse_2b	61
20	Rat_2b	61
	Rat_2a	61
	Dog_2a	61
	Cat_2a	61
	Pig_2a	61
25	Pig_2b	61
	Human_2b	61
	Human_2c	61
	Human_2a	61
	Rabbit_2a	61
30	Rabbit_2b	61
	Mouse_2a	121	EYEPENGKVYRQDRKSVQRIKAKDIVPGDIVEIAVGDKVPADIRLTSIKSTTLRVDQSIL
	Mouse_2b	121
	Rat_2b	121
35	Rat_2a	121
	Dog_2a	121
	Cat_2a	121
	Pig_2a	121
	Pig_2b	121
40	Human_2b	121
	Human_2c	121
	Human_2a	121
	Rabbit_2a	121
	Rabbit_2b	121
45	Mouse_2a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAMGVVVATGVNTEIGKIRDEM
	Mouse_2b	181
	Rat_2b	181
	Rat_2a	181
50	Dog_2a	181
	Cat_2a	181
	Pig_2a	181
	Pig_2b	181
	Human_2b	181
55	Human_2c	181
	Human_2a	181
	Rabbit_2a	181
	Rabbit_2b	181

	Mouse_2a	241	ATEQERTPLQOKLDEFGEQLSKVISLICIAVWIINIGHFNDFVHGGSWIRGAIYYFKIAV
	Mouse_2b	241
	Rat_2b	241
5	Rat_2a	241
	Dog_2a	241
	Cat_2a	241
	Pig_2a	241
	Pig_2b	241
10	Human_2b	241
	Human_2c	241
	Human_2a	241
	Rabbit_2a	241
	Rabbit_2b	241
15	Mouse_2a	301	ALAVAAIPEGLPAVITTCALGTRRMAKKNNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Mouse_2b	301
	Rat_2b	301
	Rat_2a	301
20	Dog_2a	301
	Cat_2a	301
	Pig_2a	301
	Pig_2b	301
	Human_2b	301
25	Human_2c	301
	Human_2a	301
	Rabbit_2a	301
	Rabbit_2b	301
30	Mouse_2a	361	MSVCRMFI LDKVEGDTCSLNEFSITGSTYAPIGEVQKDDKPKVCHQYDGLVELATICALC
	Mouse_2b	361
	Rat_2b	361T.....
	Rat_2a	361T.....
	Dog_2a	361R.....S.....T.....H.....
35	Cat_2a	361T.....H.....
	Pig_2a	361T.....H.....
	Pig_2b	361T.....H.....
	Human_2b	361R.....T.....H.....N.....
	Human_2c	361R.....T.....H.....N.....
40	Human_2a	361R.....T.....H.....N.....
	Rabbit_2a	361D.....T.....H.....
	Rabbit_2b	361D.....T.....H.....
45	Mouse_2a	421	NDSALDYNEAKGVYKVG EATETALTCLVEKMNVFDTELKGLSKIERANACNSVIKQLMK
	Mouse_2b	421
	Rat_2b	421
	Rat_2a	421
	Dog_2a	421
	Cat_2a	421K.F.....
50	Pig_2a	421
	Pig_2b	421
	Human_2b	421
	Human_2c	421
	Human_2a	421
55	Rabbit_2a	421
	Rabbit_2b	421

	Mouse_2a	481	KEFTLEFSRDRKSM SVYCTPNKPSRTSMSKMFVKGAPEGVIDRCTHIRVGSTKVPMTPGV
	Mouse_2b	481
	Rat_2b	481
	Rat_2a	481
5	Dog_2a	481
	Cat_2a	481
	Pig_2a	481
	Pig_2b	481
	Human_2b	481S..
10	Human_2c	481S..
	Human_2a	481S..
	Rabbit_2a	481A..
	Rabbit_2b	481A..
15	Mouse_2a	541	KQKIMSVIREWGS GSDTLRCLALATHDNPLKREEMHLEDSANFIKYETNLTFVGCVGMLD
	Mouse_2b	541
	Rat_2b	541R.....
	Rat_2a	541R.....
	Dog_2a	541	...V.....R...N.....
20	Cat_2a	541	...V.....R...N.....
	Pig_2a	541MR...N.....
	Pig_2b	541MR...N.....
	Human_2b	541R.....
	Human_2c	541R.....
25	Human_2a	541R.....
	Rabbit_2a	541R.....K.....
	Rabbit_2b	541R.....K.....
30	Mouse_2a	601	PPRIEVASSVKLC RQAGIRVIMITGDNKGTAVAICRRIGIFGQDEDVTSKAF TGREFDEL
	Mouse_2b	601
	Rat_2b	601
	Rat_2a	601
	Dog_2a	601
	Cat_2a	601
35	Pig_2a	601
	Pig_2b	601
	Human_2b	601
	Human_2c	601
	Human_2a	601
40	Rabbit_2a	601E...A.....
	Rabbit_2b	601E...A.....
	Mouse_2a	661	SPSAQRDA CLNARCFARVEPSHKSKIVEFLQSFDEITAMTGDGVNDAPALKKSEIGIAMG
	Mouse_2b	661
45	Rat_2b	661
	Rat_2a	661
	Dog_2a	661
	Cat_2a	661
	Pig_2a	661	N....E.....
50	Pig_2b	661	N....E.....
	Human_2b	661	N.....A.....
	Human_2c	661	N.....A.....
	Human_2a	661	N.....A.....
	Rabbit_2a	661	N.....A.....
55	Rabbit_2b	661	N.....A.....

	Mouse_2a	721	SGTAVAKTASEMVLADDNFSTIVAAVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAAL
	Mouse_2b	721
	Rat_2b	721
	Rat_2a	721
5	Dog_2a	721
	Cat_2a	721
	Pig_2a	721
	Pig_2b	721
	Human_2b	721
10	Human_2c	721
	Human_2a	721
	Rabbit_2a	721
	Rabbit_2b	721
15	Mouse_2a	781	GFPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMNKPPRNPKEPLISGWLFFRYLAIG
	Mouse_2b	781
	Rat_2b	781
	Rat_2a	781
	Dog_2a	781
20	Cat_2a	781
	Pig_2a	781
	Pig_2b	781
	Human_2b	781
	Human_2c	781
25	Human_2a	781
	Rabbit_2a	781
	Rabbit_2b	781
30	Mouse_2a	841	CYVGAATVGAAAWWFIAADGGPRVSFYQLSHFLQCKEDNPFDGVDCAIFESPYPMTMAL
	Mouse_2b	841
	Rat_2b	841E.....
	Rat_2a	841E.....
	Dog_2a	841D.....E.....
	Cat_2a	841D.....E.....
35	Pig_2a	841T.....E.....V.....
	Pig_2b	841T.....E.....V.....
	Human_2b	841E.....
	Human_2c	841E.....
	Human_2a	841E.....
40	Rabbit_2a	841E.....
	Rabbit_2b	841E.....
45	Mouse_2a	901	SVLVTIEMCNALNSLSENQSLLRMPWENIWLVGSIKLSMSLHFLILYVEPLPLIFQITP
	Mouse_2b	901
	Rat_2b	901
	Rat_2a	901
	Dog_2a	901
	Cat_2a	901
	Pig_2a	901
50	Pig_2b	901
	Human_2b	901
	Human_2c	901
	Human_2a	901
	Rabbit_2a	901
55	Rabbit_2b	901

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Mouse_2a  961  LNLTQWLMVLKISLPVILMDETLKFVARNYLEQPAILE-----
Mouse_2b  961  .....GKECVQPATKSSCSLSACTDGISWPF
Rat_2b    961  .....GKECA.....P.....
Rat_2a    961  .....AILE
5  Dog_2a   961  .....AILE
   Cat_2a   961  .....AILE
   Pig_2a   961  .....AILE
   Pig_2b   961  .....GKEC.....F.....
10  Human_2b 961  ..V.....GKEC.....F.....
   Human_2c 961  ..V.....VLSSL
   Human_2a 961  ..V.....AILE
   Rabbit_2a 961  ..V.....AILE
   Rabbit_2b 961  ..V.....GKEC....PQ.-...W...E.V....

15  Mouse_2b 1021  VLLIMPLVVVVYSTDTNFSDFWS
   Rat_2b   1020  .....
   Pig_2b   1019  .....
   Human_2b 1019  .....
   Rabbit_2b 1019  ....V...M.....LL..
20

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Fig. 3 Targeting construct for Serca2 flox gene modification. Sequence information.

----- Serca2 gene -----

LoxP site 1: Intron 1
 underlined sequence = loxP site and cloning sequence

Exon I (partial)
 CACGGGGCTGAGCTTGGAGCAGGTCAAGAAGCTCAAGGAGAGATGGGGCTCCAACGgtaggtgccccggc
 ccggggctgcagcggcgcgggcgggcccgagcgccaaggaagatggctgacccggctccacctcgtggg
 gcttggctcggcgcgggcgcccgacggctgcgagaggccggcggtccacgcgcgggtctgggccatcgccg
 10 accttaggggtctcgaatcaagcttatcgataccgtcgatcggacctcgagggggggcccggtacccggg
gatcaattcgagctcgcccggggatcgatccggaacccttaatATAACTTCGTATAATGTATGCTATACG
AAGTTATtaggtccctcgacctgcagcccaagctccGGGGAtctcgagccgggtgaccttcccggccggcg
 ctgagcgagtcggattgggggggggggagaggagtgaggagggagggaggttcctgcggtgggctg
 agtcccccgcgatttatgaggcgctcgatgttggtgagaaacctcgacccgtttcttggtgctcccaaa
 15 gttgcacatctggcagaagtgatgaccagctgaaatgactgcatggctcctggaggccggagagggtta
 cgggcagttccgaggccactgattaccagggtgaataattttctcggggtatcaaagtggagacagatt
 gttgtacgttcatacacctatatccgccattcagacaacgatgggtgggtgaatttagcagtttttaataaa
 agcgctaataacaatatcttcatttttctttc

----- Serca2 gene -----

LoxP site 2: Intron 3 5' of genomic XcmI site
 underlined sequence = loxP site, cloning sites and partial HSV-TK

ccaatttttattcttagaacattgtattcttatactgtgtaggaagtgaataatcatatagctacttgtc
 ttaggtttcacaaaactgataactgtatggtttcaattatgtattcacacgtttaagtctgacccagggG
 20 GATCCggaacccttaatATAACTTCGTATAATGTATGCTATACGAAGTTATtaggtccctcgacctgcag
cccaagctgatcctctagtcgagccccagctgggtcttttcgcctcagaagccatagagcccaccgcac
ccagcatgctgctattgtcttcccaatcctcccccttgctgtcctgccccacccccccagaata
gaatgacacctactcagacaatgcatgcaatttctcattttatttaggaaaggacagtgggagtggtgac
cttcagggtcgaaggaaggcagggggaggggcaaacacagatggctggcaactagaaggcacagtcga
ggctgatcagcgctctagctagagaattgatccctcagaagaactcgtcaagaaggcagtagaaggc
 30 gatgcgctgcgaatcgggagcggcgat*cggtaaagcacgaggaagcgg*cagcccatctgcgcgcaagc
tctttcagcaatatcacgggtagccaacgctatgt*ctgataagcgggtccgcccacacccaa*cggcca
caagtc*atgaaatcca*aaaaagcggggccatttttccacc*atgatttt*cggcaagcaaggccttt*c
cattgggtcaccgac*aga*catt*tccgt*c*ggcattgcgc*ccct

----- HSV-TK Neo antibiotics cassette -----

LoxP site 3: Intron 3 5' of genomic XcmI site
 underlined sequence = loxP site, cloning sites and partial Neo gene

gttttcat*accacgcgggtcccgggc*gatat*ttcaccttgtc*ag*cggtgttggtggtgtaaatg
 ttccgcatgtttcgaaagccc*agcaccgcagtaagtcacatcggtcgggtacgtagacgatatcgtc
 gcgcgaaccagggccaccagcaagttgcgtgggtgggttttcccatcc*gtggggac*gtctatataa
 40 acc*gcagtagcgtgggcattttctgctccgggcggacttccgtgggttcttgctgcccggcgagggcgcaa
cggcgtacgtcggttgctatggccgcgagaaacgcgcagcctgggtcgaacgcagacgcgtgttgatggccgg
ggtacgaagccatacgcgcttctacaaggcgctggccgaagaggtgccccgaggtttcacgccaccaagatct
gcggcacgctgttgacgctgttaagcgggtcgctgcagggtcgctcggtgttcgaggccacacgcgtcacc
ttaatatgcgaagtggacctcggaaccgcgcgccccgactgcattctgcgtgttcgaattcgccaatgacaa
 45 gacgctggggcggggtttgctcgacattgggtggaaacattccaggcctgggtggagaggctttttgcttcc
tcttgcaaaaccacactgctcgacattgggtggaaacattccaggcctgggtggagaggctttttgcttcc
tcttgaaaaccacactgctcgatccggaacccttaatATAACTTCGTATAATGTATGCTATACGAA
GTTATtaggtccctcgacctgcagcccaagctgatcctctagagtcgacctcgatctgtggtcatggcctctatgaaa
acattagcttagagg

Fig. 4A **Schematic representation of genetic manipulation.**

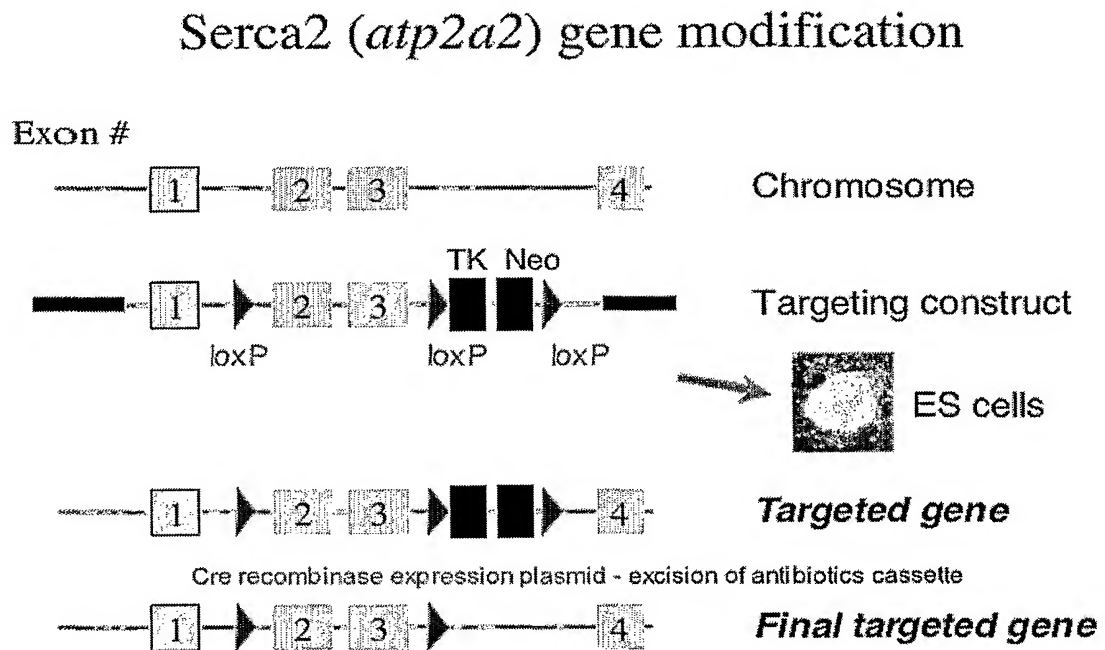


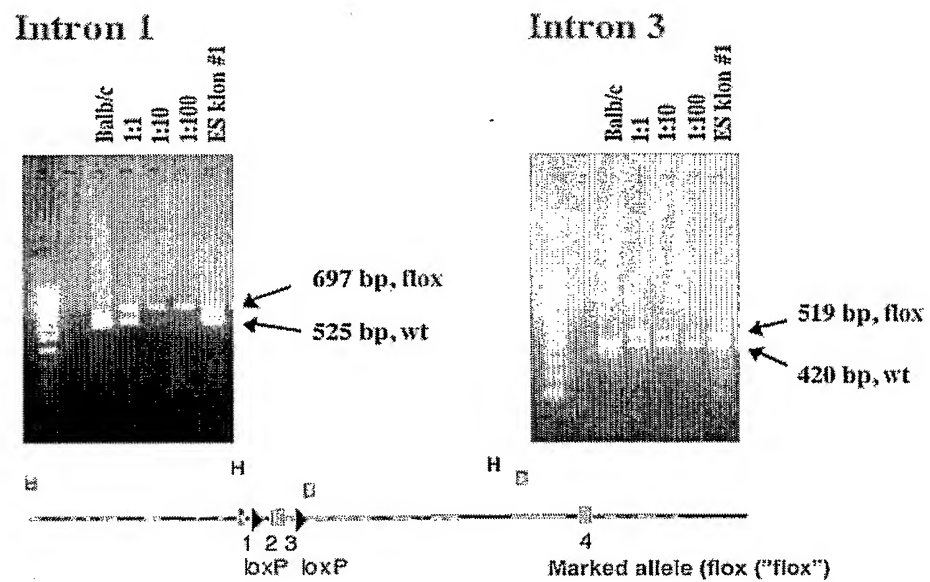
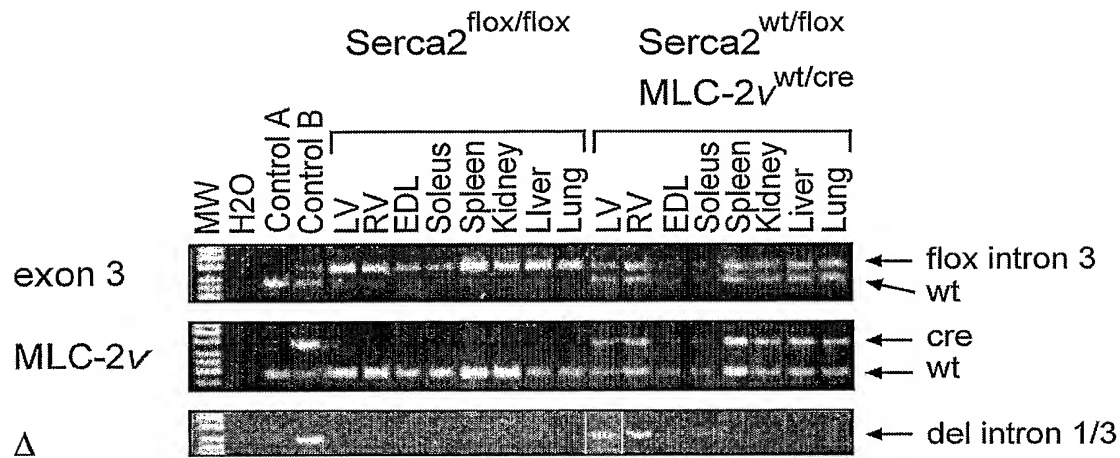
Fig 4 B: Verification of Serca locus targeting events offspring from chimeric mice.

Fig. 5 Specificity of gene deletion in a test model.

5



- 10 Control A = wt ES cells (14.1a)
 Control B = Upper panel: Serca2^{wt/flox} ES cells
 Middle panel: Left ventricle from MLC-2V-Cre mice
 Lower panel: Serca2^{wt/del} ES cells
- 15 LV = heart left ventricle
 RV = heart right ventricle
 EDL = extensor digitorum longus muscle (fast-twitch skeletal muscle)
 Soleus = soleus muscle (slow-twitch skeletal muscle)
 Other tissues as indicated.

Fig. 6 Cardiac ANP mRNA expression.

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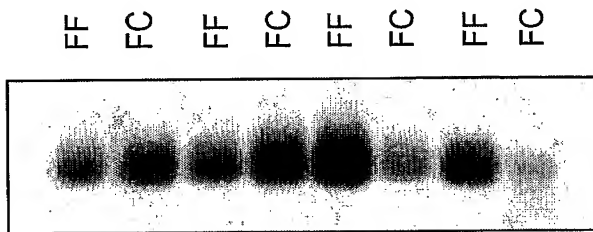
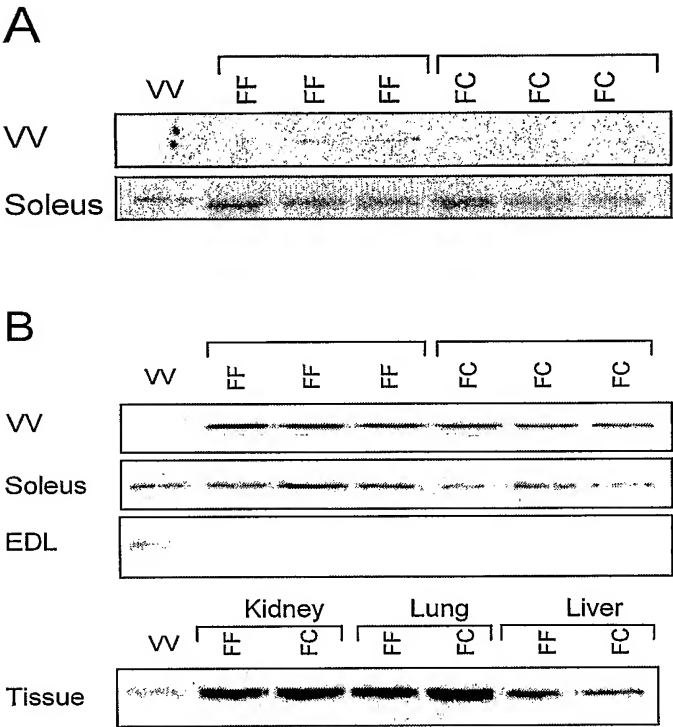


Fig. 7 Serca2 protein expression.

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Fig. 8 Compensatory mechanisms in *Serca^{fllox}* MLC-2 ν -Cre mice.

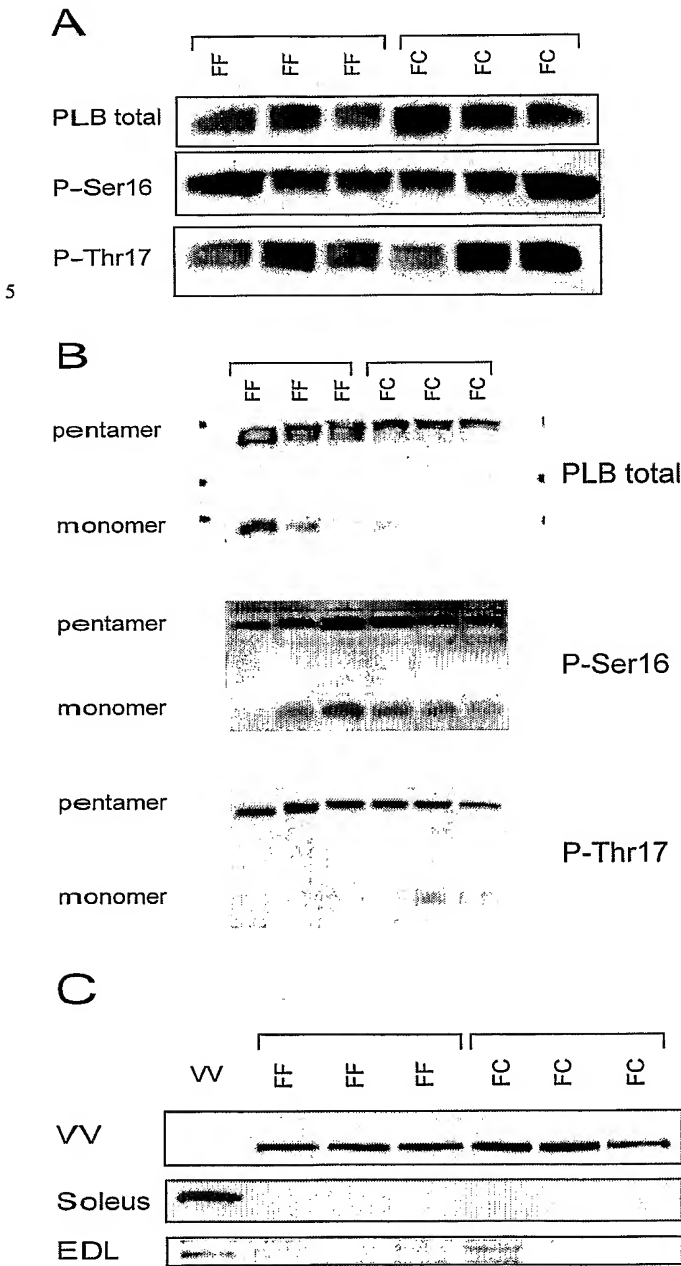
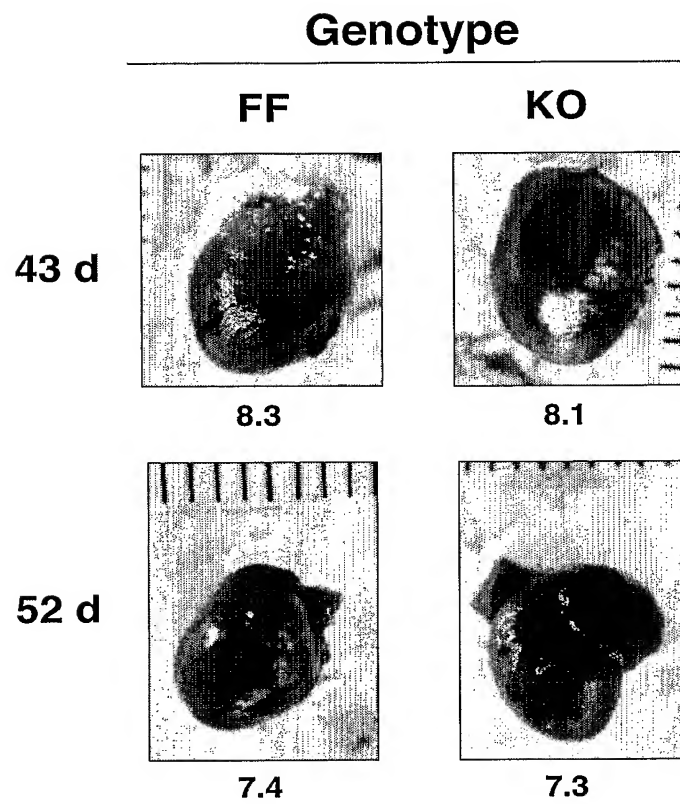


Figure 10

Heart morphology

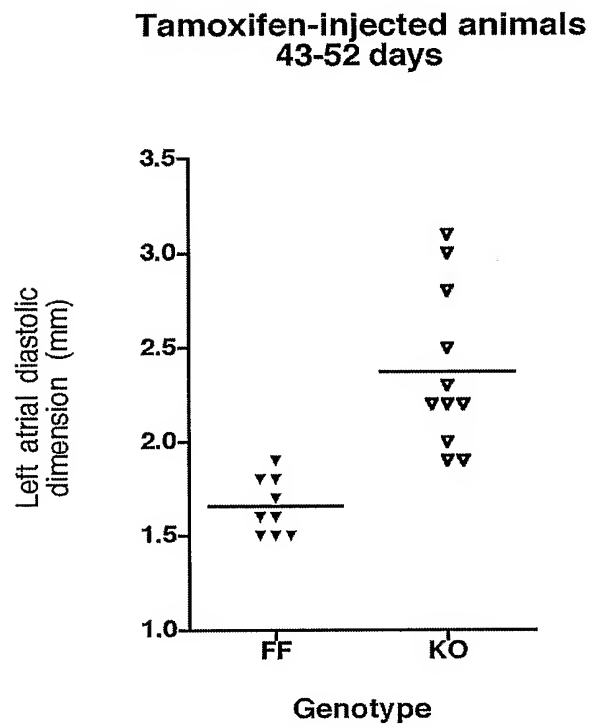
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Figure 11 Pilot series left atrial diastolic diameter.

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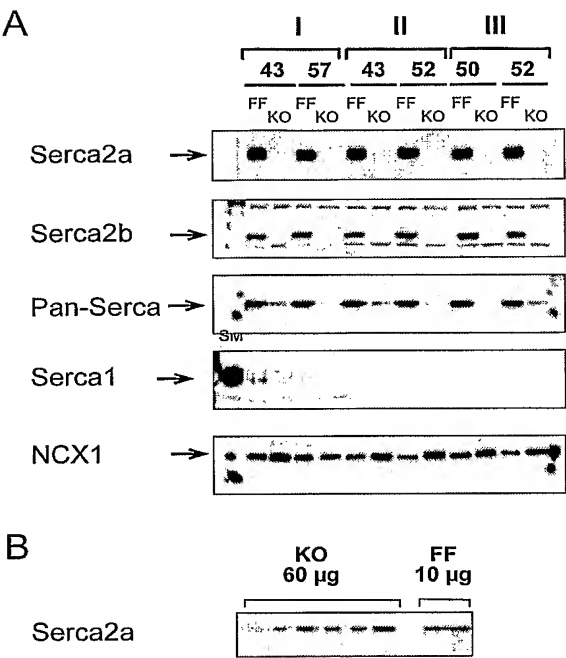


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Figure 12 Serca protein content in tamoxifen-induced FF and KO mice

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